

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/368,776DATE: 01/28/95
TIME: 16:26:27

INPUT SET: S2076.raw

This Raw Listing contains the General
Information Section and up to the first 5 pages.

ENTERED

SEQUENCE LISTING

(1) General Information:

(i) APPLICANT: Thomas Ciossek, Axel Ullrich, Birgit Millauer

(ii) TITLE OF INVENTION: METHODS FOR DIAGNOSIS AND TREATMENT OF MDK1 SIGNAL TRANSDUCTION DISORDERS

(iii) NUMBER OF SEQUENCES: 10

(iv) CORRESPONDENCE ADDRESS:

(A) ADDRESSEE: Lyon & Lyon
(B) STREET: 633 West Fifth Street
(C) CITY: Los Angeles
(D) STATE: California
(E) COUNTRY: USA
(F) ZIP: 90071

(v) COMPUTER READABLE FORM:

(A) MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
(B) COMPUTER: IBM compatible
(C) OPERATING SYSTEM: IBM P.C. DOS (Version 5.0)
(D) SOFTWARE: WordPerfect (Version 5.1)

(vi) CURRENT APPLICATION DATA:

(A) APPLICATION NUMBER: to be assigned
(B) FILING DATE: January 3, 1995
(C) CLASSIFICATION:

(vii) PRIOR APPLICATION DATA:

Prior applications total,
including application
described below: none(A) APPLICATION NUMBER:
(B) FILING DATE:

(viii) ATTORNEY/AGENT INFORMATION:

RAW SEQUENCE LISTING
PATENT APPLICATION US/08/368,776DATE: 01/28/95
TIME: 16:26:30

INPUT SET: S2076.raw

47 (A) NAME: Warburg, Richard J;
48 (B) REGISTRATION NUMBER: 32,327
49 (C) REFERENCE/DOCKET NUMBER: 208/007
50

(ix) TELECOMMUNICATION INFORMATION:

51
52
53 (A) TELEPHONE: (213) 489-1600
54 (B) TELEFAX: (213) 955-0440
55 (C) TELEX: 67-3510
56
57

(1) INFORMATION FOR SEQUENCE ID NO: 1:

(i) SEQUENCE CHARACTERISTICS:

60
61 (A) LENGTH: 4304
62 (B) TYPE: nucleic acid
63 (C) STRANDNESS: single
64 (D) TOPOLOGY: linear
65

(ii) MOLECULE TYPE: nucleic

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

66
67
68
69
70
71 AAGCGGCCGG TCTGCAGTCG GAGACTTGCA GGCAGCAAAC ACGGTGCGAA 50
72
73 CGAACCGGAG GGGGGAGAGA GAAATCAAAC AGCTAAGCGT GGAGCAGACG 100
74
75 GCCTGGGACC CAGAAGGGGA TCGATGCGAG GAGCGCAATA ATAACAACAA 150
76
77 TAATAACCCA CTTCGGAGCA AACAGCATCT AAAGAGCTGC GACCCAAC TG 200
78
79 CAGCCTAAAA AAATCAAACC TGCTCATGCA CCATGGTTGT TCAAAC TCGG 250
80
81 TTCCCTTCGT GGATTATTTT GTGTTACATC TGGCTGCTTG GCTTTGCACA 300
82
83 CACGGGGGAG GCGCAGGCTG CGAAGGAAGT ACTATTACTG GACTCGAAAG 350
84
85 CACAACAAAC AGAATTGGAA TGGATTTCCCT CTCCACCCAG TGGGTGGGAA 400
86
87 GAAATTAGTG GTTTGGATGA GAACTACACT CCGATAAGAA CATAACAGGT 450
88
89 GTGCCAGGTC ATGGAGCCCA ACCAGAACAA CTGGCTGCGG ACTAACTGGA 500
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91 TTTCTAAAGG CAACGCACAA AGGATTTTTG TAGAATTGAA ATTCACCTTG 550
92
93 AGGGATTGTA ATAGTCTTCC CGGAGTCCTG GGAAC TTGCA AGGAAACGTT 600
94
95 TAATTTGTAC TATTATGAAA CAGACTACGA CACCGGCAGG AATATACGAG 650
96
97 AAAACCTTTA TGTTAAAATA GACACCATTG CTGCAGATGA AAGTTTCACA 700
98
99 CAAGGTGACC TTGGTGAAAG AAAGATGAAG CTGAACACTG AGGTGAGAGA 750

RAW SEQUENCE LISTING
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TIME: 16:26:33

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149	GTTCGGCTTC	ATCATTTGGAA	GAAGGCACTG	TGGTTATAGC	AAGGCTGACC	2000
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RAW SEQUENCE LISTING
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165	GTCATGATTG	TGATAGAGTT	CATGGAGAAT	GGAGCCCTGG	ATGCATTTCT	2400
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167	CAGGAAACAC	GATGGGCAGT	TTACAGTCAT	TCAGTTGGTA	GGAATGTTGA	2450
168						
169	GAGGTATTGC	CGCTGGGATG	CGATACTTGG	CTGATATGGG	ATACGTTCAC	2500
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171	AGGGACCTTG	CAGCGCGCAA	CATCCTTGTC	AACAGCAATC	TTGTTTGTA	2550
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173	AGTGTCAGAT	TTTGGCCTTT	CCCGGGTTAT	AGAGGATGAT	CCCGAAGCTG	2600
174						
175	TCTACACCAC	GACTGGTGGA	AAAATTCCAG	TAAGGTGGAC	TGCACCGGAA	2650
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177	GCCATTCAAT	ACCGGAAGTT	CACCTCAGCC	AGCGATGTGT	GGAGCTATGG	2700
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183	GCGCCCATGG	ATTGCCCAGC	TGGTCTTCAC	CAGCTAATGC	TGGATTGTTG	2850
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187	TAGACAAAAT	GATTCGAAAC	CCAAGTAGTC	TGAAAACACC	CCTGGGAACT	2950
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193	ATAAGGACAA	CTTCACAGCA	GCGGGTTACA	ACTCACTCGA	GTCAGTGGCC	3100
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195	AGGATGACTA	TCGATGATGT	GATGAGTTTA	GGGATCACAC	TGGTTGGCCA	3150
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197	TCAAAAAGAAG	ATCATGAGCA	GCATCCAGAC	TATGCGGGCA	CAAATGTTGC	3200
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199	ATTTACACGG	AACAGGCATC	CAAGTGTGAC	ACATCGGCCT	CCCTCAGATG	3250
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RAW SEQUENCE LISTING
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213	GAAAGAATAT	ATTTGTCAAA	TAAAAATCAT	GATCTTATTG	TTAAAATCAA	3600	
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215	TGAAATATTT	TCCTTAAAAT	ATGTGATTTT	AGACTATTCT	TTTCCAGAAC	3650	
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217	CATCTGTGTT	TATTCTGCTT	AAGGACTTTG	TTTTAGAAAG	TTATTTGTAG	3700	
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219	CTTTGGACCT	TTTTAGTGTT	AAATTTATGA	CACGTTACTA	CACTGGGAAC	3750	
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221	CTTTGAAGAC	TCTCAAACCT	AAAGGAAAGC	AAACTACGC	ACATAGTCGA	3800	
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223	GGATGGACTT	TGTCCTTCAT	GGCTTTGGTA	TCCTGGCTGT	GTCATTTTGT	3850	
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225	TAAACCAGTG	ATGTTTTTCAT	ATTGTTTGCT	GATTGGCAGG	TAGTTCAAAA	3900	
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231	AAGCAATAAT	GATCCATAAA	TACTATAAGG	CACTTTTAAC	AGATTGTTTA	4050	
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233	TAGAGTGATT	TACTAGGCAG	AATTTAATAA	AAAAAAAAGA	GAGATGTCAA	4100	
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235	ATTTTAGGTT	TATGTGTATA	TGATAAAAGG	CTGAGCTTCG	TCTGAAGATG	4150	
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237	CTGGTGAAAG	CAAGACTGGA	AGCGAAGCTC	TCCAGCTTTG	GCTAACCCAA	4200	
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239	TCCGAGCACA	TCAAGAGCTT	CAGTCTTGTT	ACAGTAAGAA	ATTTAGGAAC	4250	
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243	AAAA					4304	
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245							
246							

(1) INFORMATION FOR SEQUENCE ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

(A)	LENGTH:	998
(B)	TYPE:	amino acid
(C)	STRANDNESS:	single
(D)	TOPOLOGY:	linear

(ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

PAGE: 1

SEQUENCE VERIFICATION REPORT
PATENT APPLICATION US/08/368,776

DATE: 01/28/95
TIME: 16:26:44

INPUT SET: S2076.raw

Line	Error	Original Text
32	Wrong application Serial Number	(A) APPLICATION NUMBER: to be assigned